

NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 CFR 1.821 - 1.825 for the following reason(s):

- ☐ 1. This application clearly fails to comply with the requirements of 37 CFR 1.821 - 1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.
- ☐ 2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 CFR 1.821(c).
- ☒ 3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 CFR 1.821(e).
- ☐ 4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 CFR 1.822 and/or 1.823, as indicated on the attached marked-up copy of the "Raw Sequence Listing."
- ☐ 5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A substitute computer readable form must be submitted as required by 37 CFR 1.825(d).
- ☐ 6. The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 CFR 1.821(e).
- ☐ 7. Other: _____

Applicant must provide:

- ☐ An initial or substitute computer readable form (CRF) copy of the "Sequence Listing"
- ☐ An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification
- ☒ A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 CFR 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d)

For questions regarding compliance with these requirements, please contact:

For Rules Interpretation, call (703) 308-1123
For CRF submission help, call (703) 308-4212
For PatentIn software help, call (703) 308-6856

Please return a copy of this notice with your response.

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/487,461DATE: 09/26/95
TIME: 15:55:05

INPUT SET: S6366.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

SEQUENCE LISTING

(1) General Information:

- (i) APPLICANT: Parker, Thomas S.; Levine, Daniel M.;
Rubin, Albert L; Gordon, Bruce R; Saal, Stuart D.
- (ii) TITLE OF INVENTION: Methods Useful in Endotoxin Based
Prophylaxis and Therapy
- (iii) NUMBER OF SEQUENCES: 9
- (iv) CORRESPONDENCE ADDRESS:
(A) ADDRESSEE: Felfe & Lynch
(B) STREET: 805 Third Avenue
(C) CITY: New York City
(D) STATE: New York
(E) COUNTRY: USA
(F) ZIP: 10022
- (v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE: Diskette, 5.25 inch, 360 kb
storage
(B) COMPUTER: IBM PS/2
(C) OPERATING SYSTEM: PC-DOS
(D) SOFTWARE: Wordperfect
- (vi) CURRENT APPLICATION DATA:
(A) APPLICATION NUMBER: 08/487,461
(B) FILING DATE: June 7, 1995
(C) CLASSIFICATION: 435
- (vii) PRIOR APPLICATION DATA:
(A) APPLICATION NUMBER: PCT/US93/07453
(B) FILING DATE: 9-AUGUST-1993
- (vii) PRIOR APPLICATION DATA:
(A) APPLICATION NUMBER: 07/928,930
(B) FILING DATE: 12-AUGUST-1992
- (viii) ATTORNEY/AGENT INFORMATION:
(A) NAME: Hanson, Norman D.
(B) REGISTRATION NUMBER: 30,946
(C) REFERENCE/DOCKET NUMBER: ROGO 211.2
- (ix) TELECOMMUNICATION INFORMATION:

ENTERED

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RAW SEQUENCE LISTING
PATENT APPLICATION US/08/487,461DATE: 09/26/95
TIME: 15:55:08

INPUT SET: S6366.raw

47 (A) TELEPHONE: (212) 688-9200
48 (B) TELEFAX: (212) 838-3884
49
50 (2) INFORMATION FOR SEQ ID NO: 1:
51 (i) SEQUENCE CHARACTERISTICS:
52 (A) LENGTH: 18 amino acids
53 (B) TYPE: amino acid
54 (D) TOPOLOGY: linear
55 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
56
57 Asp Trp Leu Lys Ala Phe Tyr Asp Lys Val Ala Glu Lys Leu Lys
58 5 10 15
59 Glu Ala Phe
60
61
62
63
64 (2) INFORMATION FOR SEQ ID NO: 2:
65 (i) SEQUENCE CHARACTERISTICS:
66 (A) LENGTH: 18 amino acids
67 (B) TYPE: amino acid
68 (D) TOPOLOGY: linear
69 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
70
71 Lys Trp Leu Asp Ala Phe Tyr Lys Asp Val Ala Lys Glu Leu Glu
72 5 10 15
73 Lys Ala Phe
74
75
76
77
78 (2) INFORMATION FOR SEQ ID NO: 3:
79 (i) SEQUENCE CHARACTERISTICS:
80 (A) LENGTH: 17 amino acids
81 (B) TYPE: amino acid
82 (D) TOPOLOGY: linear
83 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
84
85
86 Asp Trp Leu Lys Ala Phe Tyr Asp Lys Ala Glu Lys Leu Lys Glu
87 5 10 15
88 Ala Phe
89
90
91
92
93 (2) INFORMATION FOR SEQ ID NO: 4:
94 (i) SEQUENCE CHARACTERISTICS:
95 (A) LENGTH: 22 amino acids
96 (B) TYPE: amino acid
97 (D) TOPOLOGY: linear
98 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
99

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/487,461DATE: 09/26/95
TIME: 15:55:11

INPUT SET: S6366.raw

100 Pro Lys Leu Glu Glu Leu Lys Glu Lys Leu Lys Glu Leu Leu Glu
101 5 10 15

102 Lys Leu Lys Glu Lys Leu Ala
103 20

104

105

106

107 (2) INFORMATION FOR SEQ ID NO: 5:

108 (i) SEQUENCE CHARACTERISTICS:

109 (A) LENGTH: 16 amino acids

110 (B) TYPE: amino acid

111 (D) TOPOLOGY: linear

112 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

113

114 Val Ser Ser Leu Lys Glu Tyr Trp Ser Ser Leu Lys Glu Ser Phe
115 5 10 15

116 Ser

117

118

119

120 (2) INFORMATION FOR SEQ ID NO: 6:

121 (i) SEQUENCE CHARACTERISTICS:

122 (A) LENGTH: 20 amino acids

123 (B) TYPE: amino acid

124 (D) TOPOLOGY: linear

125 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

126

127 Val Ser Ser Leu Leu Ser Ser Leu Lys Glu Tyr Trp Ser Ser Leu
128 5 10 15

129 Lys Glu Ser Leu Ser

130 20

131

132

133 (2) INFORMATION FOR SEQ ID NO: 7:

134 (i) SEQUENCE CHARACTERISTICS:

135 (A) LENGTH: 24 amino acids

136 (B) TYPE: amino acid

137 (D) TOPOLOGY: linear

138 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

139

140 Val Ser Ser Leu Leu Ser Ser Leu Leu Ser Ser Leu Lys Glu Tyr
141 5 10 15

142 Trp Ser Ser Leu Lys Glu Ser Glu Ser

143 20

144

145

146

147 (2) INFORMATION FOR SEQ ID NO: 8:

148 (i) SEQUENCE CHARACTERISTICS:

149 (A) LENGTH: 22 amino acids

150 (B) TYPE: amino acid

151 (D) TOPOLOGY: linear

152 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/487,461DATE: 09/26/95
TIME: 15:55:13

INPUT SET: S6366.raw

153
154 Pro Val Leu Asp Glu Phe Arg Glu Lys Leu Asn Glu Glu Leu Glu
155 5 10 15
156 Ala Leu Lys Gln Lys Met Lys
157 20
158
159
160
161 (2) INFORMATION FOR SEQ ID NO: 9:
162 (i) SEQUENCE CHARACTERISTICS:
163 (A) LENGTH: 22 amino acids
164 (B) TYPE: amino acid
165 (D) TOPOLOGY: linear
166 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
167
168 Pro Leu Ala Glu Asp Leu Gln Thr Lys Leu Asn Glu Asn Val Glu
169 5 10 15
170 Asp Leu Arg Lys Gln Leu Val
171 20
172